

F Errors Corrected by the STIC Systems Branch

Serial Number: 08/202,526

CRF Processing Date: 1/9/98
 Edited by: hw
 Verified by: hw (STIC staff)

ENTERED

#10.5

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☒ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

***Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.**

P. Gambel

1806

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/702,525

DATE: 01/09/98
TIME: 14:06:23

INPUT SET: S22465.raw

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

Does Not Comply
Corrected Diskette Needed

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SEQUENCE LISTING

(1) General Information:

(i) APPLICANTS: Sharpe, Sharpe
Borriello, Francescopaolo
Freeman, Gordon
Nadler, Lee

(ii) TITLE OF INVENTION: Novel Forms of T Cell Costimulatory Molecules
and Uses Therefor

(iii) NUMBER OF SEQUENCES: 65

(iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: LAHIVE & COCKFIELD
(B) STREET: 28 State Street
(C) CITY: Boston
(D) STATE: Massachusetts
(E) COUNTRY: USA
(F) ZIP: 02109-1875

(v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Floppy disk
(B) COMPUTER: IBM PC compatible
(C) OPERATING SYSTEM: PC-DOS/MS-DOS
(D) SOFTWARE: ASCII Text

(vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:

(vi) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER: US 08/205,697
(B) FILING DATE: 02-Mar-1994

(viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Mandragouras, Amy E.
(B) REGISTRATION NUMBER: 36,207
(C) REFERENCE/DOCKET NUMBER: BWI-120CPUS

(ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: (617)227-7400
(B) TELEFAX: (617)227-5941

INPUT SET: S22465.raw

46
47

ERRORED SEQUENCES FOLLOW:

3001 (2) INFORMATION FOR SEQ ID NO:65:
3002
3003 (i) SEQUENCE CHARACTERISTICS:
3004 (A) LENGTH: 226 amino acids
3005 (B) TYPE: amino acid
3006 (D) TOPOLOGY: linear
3007
3008 (ii) MOLECULE TYPE: protein
3009
3010 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:
3011
3012 Met Ala Cys Asn Cys Gln Leu Met Gln Asp Thr Pro Leu Leu Lys Phe
3013 1 5 10 15
3014
3015 Pro Cys Pro Arg Leu Ile Leu Leu Phe Val Leu Leu Ile Arg Leu Ser
3016 20 25 30
3017
3018 Gln Val Ser Ser Asp Val Asp Glu Gln Leu Ser Lys Ser Val Lys Asp
3019 35 40 45
3020
3021 Lys Val Leu Leu Pro Cys Arg Tyr Asn Ser Pro His Glu Asp Glu Ser
3022 50 55 60
3023
3024 Glu Asp Arg Ile Tyr Trp Gln Lys His Asp Lys Val Val Leu Ser Val
3025 65 70 75 80
3026
3027 Ile Ala Gly Lys Leu Lys Val Trp Pro Glu Tyr Lys Asn Arg Thr Leu
3028 85 90 95
3029
3030 Tyr Asp Asn Thr Thr Tyr Ser Leu Ile Ile Leu Gly Leu Val Leu Ser
3031 100 105 110
3032
3033 Asp Arg Gly Thr Tyr Ser Cys Val Val Gln Lys Lys Glu Arg Gly Thr
3034 115 120 125
3035
3036 Tyr Glu Val Lys His Leu Ala Leu Val Lys Leu Ser Ile Lys Pro Pro
3037 130 135 140
3038
3039 Glu Asp Pro Pro Asp Ser Lys Asn Thr Leu Val Leu Phe Gly Ala Gly
3040 145 150 155 160
3041
3042 Phe Gly Ala Val Ile Thr Val Val Val Ile Val Val Ile Ile Lys Cys
3043 165 170 175
3044
3045 Phe Cys Lys His Gly Leu Ile Tyr His Leu Gln Leu Thr Ser Ser Ala

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/702,525DATE: 01/09/98
TIME: 14:06:30

INPUT SET: S22465.raw

3046		180		185		190
3047						
3048	Lys Asp Phe Arg Asn Leu Ala Leu Pro Trp Leu Cys Lys His Gly Ser					
3049	195		200		205	
3050						
3051	Leu Gly Glu Ala Ser Ala Val Ile Cys Arg Ser Thr Gln Thr Asn Glu					
3052	210		215		220	
3053						
3054	Pro Gln					
3055	225					
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--> 3057	{PAGE 99}					
3058						
3059						
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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/702,525

DATE: 01/09/98
TIME: 14:06:32

INPUT SET: S22465.raw

Line	Error	Original Text
3057	Wrong Amino Acid Designator	{PAGE 99}